



SEQUENCE LISTING

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Zhang, Renhuai

<120> Recombinant Super-Compound Interferon

<130> 792-A-PCT-US

<140> 10/650,365
<141> 2003-08-28

<150> PCT/CN02/00128
<151> 2002-02-28

<150> CN 01104367.9
<151> 2001-02-28

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<170> PatentIn version 3.2

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atc ctg ctg gct cag atg cgt cgt atc tcc ccg ttc tcc tgc ctg aaa 96
Ile Leu Leu Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys
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gac cgt cac gac ttc ggt ttc ccg cag gaa gaa ttc gac ggt aac cag 144
Asp Arg His Asp Phe Gly Phe Pro Gln Glu Glu Phe Asp Gly Asn Gln
35 40 45
ttc cag aaa gct cag gct atc tcc gtt ctg cac gaa atg atc cag cag 192
Phe Gln Lys Ala Gln Ala Ile Ser Val Leu His Glu Met Ile Gln Gln
50 55 60
acc ttc aac ctg ttc tcc acc aaa gac tcc tcc gct gct tgg gac gaa 240
Thr Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu
65 70 75 80
tcc ctg ctg gaa aaa ttc tac acc gaa ctg tac cag cag ctg aac gac 288
Ser Leu Leu Glu Lys Phe Tyr Thr Glu Leu Tyr Gln Gln Leu Asn Asp
85 90 95

ctg gaa gct tgc gtt atc cag gaa gtt ggt gtt gaa gaa acc ccg ctg	336
Leu Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Thr Pro Leu	
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Met Asn Val Asp Ser Ile Leu Ala Val Lys Lys Tyr Phe Gln Arg Ile	
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Thr Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val	
130 135 140	

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Val Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln	
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Phe Gln Lys Ala Gln Ala Ile Ser Val Leu His Glu Met Ile Gln Gln
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Thr Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu
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Ser Leu Leu Glu Lys Phe Tyr Thr Glu Leu Tyr Gln Gln Leu Asn Asp
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Leu Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Thr Pro Leu
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Met Asn Val Asp Ser Ile Leu Ala Val Lys Lys Tyr Phe Gln Arg Ile
 115 120 125

Thr Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val
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Glu Arg Leu Arg Arg Lys Glu
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 Ile Leu Leu Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys
 20 25 30
 gac cgt cac gac ttc ggc ttt ccg caa gaa gag ttc gat ggc aac caa 144
 Asp Arg His Asp Phe Gly Phe Pro Gln Glu Glu Phe Asp Gly Asn Gln
 35 40 45
 ttc cag aaa gct cag gca atc tct gta ctg cac gaa atg atc caa cag 192
 Phe Gln Lys Ala Gln Ala Ile Ser Val Leu His Glu Met Ile Gln Gln
 50 55 60
 acc ttc aac ctg ttt tcc act aaa gac agc tct gct gct tgg gac gaa 240
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 65 70 75 80
 agc ttg ctg gag aag ttc tac act gaa ctg tat cag cag ctg aac gac 288
 Ser Leu Leu Glu Lys Phe Tyr Thr Glu Leu Tyr Gln Gln Leu Asn Asp
 85 90 95
 ctg gaa gca tgc gta atc cag gaa gtt ggt gta gaa gag act ccg ctg 336
 Leu Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Thr Pro Leu
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Ile Leu Leu Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys
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Asp Arg His Asp Phe Gly Phe Pro Gln Glu Glu Phe Asp Gly Asn Gln
 35 40 45

Phe Gln Lys Ala Gln Ala Ile Ser Val Leu His Glu Met Ile Gln Gln
 50 55 60

Thr Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu
 65 70 75 80

Ser Leu Leu Glu Lys Phe Tyr Thr Glu Leu Tyr Gln Gln Leu Asn Asp
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Leu Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Thr Pro Leu
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Met Asn Val Asp Ser Ile Leu Ala
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